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Query Match Best Local S Matches 21

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InterPro; IPR001841; Znf_ring.
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SMART; SM000064; FYVE; 1.
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PROSITE; PS0089; ZF_RING_2; 1.
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NCBI_TaxID=6239;
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use by non-profit institutions as long as its content
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STRAIN-BRISTOL N2;
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                Caldwell K.A.,
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                                                                    Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
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ZN_FING
SEQUENCE
                                                                                                                                                        Bouchard M.L., Cote S.;
"The Drosophila melanogaster developmental gene zinc-finger-motif protein.";
                                                                                                                                                                                                                                                                                     Ephydroidea; Drosophilidae; Drosophila. NCBI_TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                         006003;
01-FEB-1994 (Rel. 28, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SMART; SM00184; RING; 1.
SMART; SM00356; ZnF_C3H1; 3.
PROSITE; PS00518; ZF_RING_1;
PROSITE; PS50089; ZF_RING_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and for entities requires a license agreement (See http://www.isb-sib.
                                                                                                                                                                                                                                                                                                                                         Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Trac
Pterygota; Neoptera; Endopterygota;
                                                                                                                                                                                                                                                                                                                                                                                                      Goliath protein (G1 protein). GOL OR G1 OR GL.
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InterPro; IPR001841; Znf_ring.
Pfam; PF00097; Zf-C3HC4; 1.
Pfam; PF00642; Zf-CCCH; 3.
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      Gene 125:205-209(1993)
                                                                                                                                                                                                                    TISSUE=Embryo;
MEDLINE=93216124; PubMed=8462875;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GOLI_DROME
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"Imprinting of
                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
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-!- SIMILARITY: CONTAINS 3 C3H1-TYPE ZINC
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PUTATIVE ROLE AS TRANSCRIPTION
SUBCELLULAR LOCATION: Nuclear
TISSUE SPECIFICITY: VISCERAL ME
MUSCULATURE:
                  SIMILARITY:
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19; Conservative
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or send a
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Pfam; PF00097; zf-C3HC4; 1.
SMART; SM0184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; M97204; AAA28582.1; -. PIR; JC1495; JC1495. FlyBase; FBgn0004919; gol.
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between
                                                                                                                                                         Tsukahara F., Hattori M., Muraki T., Sakaki Y.; "Identification and cloning of a novel cDNA belonging tetratricopeptide repeat gene family from Down syndron
                                                                                                                                                                                                     TISSUE=Fetal brain; MEDLINE=97103476; P
                                                                                                                                                                                                                                                                                                                  Ohira M., Ootsuyama A., Suz
Nagase T., Nomura N., Ohki
                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation
                                                                                                                                           region 21q22.2.";
                                                                                                                                                                                                                                                                                                 Nagase T., Nomura N., Ohki M.;
"Identification of a novel human gene containing the
                                                                                                                                                                                                                                                                                                                                   MEDLINE-96281123; PubMed-8724848; Ohira M., Ootsuyama A., Suzuki E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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                                                                                                                                                                                                                                                                            tetratricopeptide repeat domain from chromosome 21.";
                                                                                                                                                                                                                                                                                                                                                                   TISSUE-Brain;
                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                             DNA Res. 3:9-16(1996).
                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
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                                                   BIOCHEM: 120:820-827(1996).
ALTERNATIVE PRODUCTS: 3 ISOFORMS; TRPDI (SHOWN HERE),
AND TPRDIII; SEEMS TO BE PRODUCED BY ALTERNATIVE SPLIC
TISSUE SPECIFICITY: FOUND IN ALL TISSUES EXAMINED.
SIMILARITY: CONTAINS 4 TPR REPEATS.
SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
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 European
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an email to license@isb-sib.ch).
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284 AA;
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Bioinformatics Institute.
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31973 MW;
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Pred. No. 0.00
8; Mismatches
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                                                                                                                                                                                                                                                                                       the Down syndrome region
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01-MAR-2002 (Rel. 41, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation updat
Makorin 4 (Zinc-finger protein 127-Xp) (ZNF
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Pfam; PF00515; TPR; 4
Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00184; RING; 1.
SMART; SM00028; TPR; 3.
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DOMAIN
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ZN_FING
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PROSITE; PS50089; ZF_RING_2; 1.
Repeat; TPR repeat; Zinc-finger;
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                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
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MKRN4 OR ZNF127L1.
                                                                                                                                                                                                                                                                      Q13434;
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REPEAT
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          the European Bioinformatics Institute. There use by non-profit institutions as long as modified and this statement is not removed. U
                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                              Hendrich B.D. Willard H.F.;
                                                                                                                                       SEQUENCE FROM N.A. Hendrich B.D., Lon
                                                                                                                                                                                                                                                                                MKR4_HUMAN
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                                                                                                                                                                       NCBI_TaxID=9606;
                                                                                                                                                                                                     Homo sapiens (Human)
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                                                                                                                 "An X-linked homologue of the autosomal imprinted
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D84294; BAA12301.1; -.
D84295; BAA12302.1; -.
D84296; BAA12303.1; -.
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17; Conser
requires a license agreement (See http://www.isb-sib.ch/announce,
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ARG/LYS-RICH (BASIC).
ARG/LYS-RICH (BASIC).
ARG/LYS-RICH (BASIC).
ARG/LYS-RICH (BASIC).
MISSING (IN ISOFORM TPRDII).
MISSING (IN ISOFORM TPRDIII).
MW; 1B4BCAA3684B6253 CRC64;
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Pred. No. 0.00
7; Mismatches
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Catarrhini;
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                                                                                                                                                                                                                                                                                PRT;
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YHBO_YEAST
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InterPro; IPRO01841; Znf_ring.
Pfam; PF00097; zf-C3HC4; 1.
Pfam; PF00642; zf-CCCH; 4.
SMART; SM00184; RING; 1.
SMART; SM00186; ZnF_C3H1; 4.
PROSITE; PS00518; ZF_RING_1; 1.
PROSITE; PS00518; ZF_RING_2; 1.
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P38748;
O1-FEB-1995
O1-FEB-1995
16-OCT-2001
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ZN_FING
ZN_FING
DOMAIN
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ZN_FING
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                                                                                                                                                                                                                               the
use
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-I- SIMILARITY: TO C.ELEGANS EEED8.9 AND S.POMBE
-I- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
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Vignati D., Wilcox L.,
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Saccharomycetales;
NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                              "Complete nucleotide sequence VIII.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomyc
Saccharomycetales; Saccharomycetaceae; Sa
                                                                                                                EMBL;
                                                                                                                                                            or send
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MEDLINE=94378003; PubMed=8091229;
                             InterPro;
                                                  InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vaudin M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A
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                                                                                                                                                                                                 s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation. European Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way iffed and this statement is not removed. Usage by and for commercial iffed and this statement is not removed. Usage by and for commercial
                           S0001002; YHL010C.
rPro; IPR001841; Znf_ring.
rPro; IPR001607; zf-UBP.
                                                                                            ; U11582; CAB34905.1; S46825; S46825.
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                                                                                                                                                            an email to license@isb-sib.ch).
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32.8%;
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protein in PRPS4-STE2
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Pred. No. 0.00
12; Mismatches
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C3H1-TYPE 2.
C3H1-TYPE 3.
MAKORIN-TYPE 6.
RING-TYPE 4.
C3H1-TYPE 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                  of Saccharomyces
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AE28B962544CEFFE CRC64;
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No. 0.00047;
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CGHTFHLQCLIQWFETAPSRT

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Similarity 20; Conser

Conservative

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RESULT 8
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Best Local
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Pfam; PF02148; zf-UBP; 1.
SMART; SM00184; RING; 1.
SMART; SM00290; ZnF_UBP; 1.
SMART; SM00290; ZnF_UBP; 1.
PROSITE; PS00518; ZF_RING_1; FALSE_NEG.
PROSITE; PS50089; ZF_RING_2; 1.
Hypothetical protein; Zinc-finger
ZN_FING 240 280 RING-TYPE.
                                                                                                                                                                                                                                                                                                                                                                    STRAIN-BRISTOL N2;
MEDLINE-92168156; PubMed-1538779;
Sulston J., Du Z., Thomas K., Wilson R., Hillier L.,
Halloran N., Green P., Thierry-Mieg J., Qiu L., Dear
Craxton M., Durbin R.K., Berks M., Metzstein M., Hawk
Ainscough R., Waterston R.;
                                                                PROSITE; PS00518; ZF_RING_1; FALSE_NEG PROSITE; PS50089; ZF_RING_2; 1.

Hypothetical protein; Zinc-finger.
Hypothetical protein; Zinc-finger.
ZN_FING 72 134

SEQUENCE 161 AA; 18847 MW; F5EF9F3;
                                                                                                                                 PIR; S15788; S15788.

wormbep; ZK637.14; CE00432.

InterPro; IPRR001841; Znf_ring

pfam; pF00097; zf-C3HC4; 1.

SMART; SM00184; RING; 1.
                                                                                                                                                                                                                            entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                     modified and this statement
                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBI the European Bioinformatics Institute. There are no restrict
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hypothetical ZK637.14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-OCT-1993
16-OCT-2001
                                                                                                                                                                                                    EMBL; Z11115; CAA77447.1;
                                                                                                                                                                                                                                                                                                                                                "The C. elegans genome sequencing project: Nature 356:37-41(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Caenorhabditis elegans.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=6239;
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27, Last sequence update)
40, Last annotation update)
kDa protein ZK637.14 in chr
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               30.5%;
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Caenorhabditis.
Score 89.5; D
Pred. No. 0.00
5; Mismatches
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                                                                   RING-TYPE.; F5EF9F3A83A9C027 CRC64;
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Pred. No. 0.00056;
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DB 1;
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nes 20;
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                                                                                                                                                                                                                                                                                                                                                            a beginning.";
                                                                                                                                                                                                                                                     Usage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19;
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                        Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                      , Staden R.,
r S., Coulson
wkins T.,
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RESULT 9
RN12_MOUSE
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                                                                       Query Match
Best Local
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16-OCT-2001 (Rel. 40, 0
16-OCT-2001 (Rel. 40, 1
16-OCT-2001 (Rel. 40, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
RING finger protein 12 (LIM domain interacting RI
(RING finger LIM domain-binding protein) (R-LIM).
RNP12 OR RLIM.
                                                                                                                               DOMAIN
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                        the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nat.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=99364422; PubMed=10431247;
Bach I., Rodriguez-Esteban C., Carriere C., Bhushan A., Krones A.,
Rose D.W., Glass C.K., Andersen B., Izpisua Belmonte J.C.,
Rosenfeld M.G.;
"Rosenfeld M.G.;
"RLIM inhibits functional activity of LIM homeodomain transcription
factors via recruitment of the histone deacetylase complex.";
Nat. Genet. 22:394-399(1999).
                                                                                                                                                                                                                                                                 EMBL; AF069992; AAD34209.1; -. MGD; MGI:1342291; Rnf12. InterPro; IPR001841; Znf_ring.
                                                                                                                                                                                                                                                                                                                          use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                      This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                               PROSITE; PS00518; ZF_RING_1; FALSE_NEG
PROSITE; PS50089; ZF_RING_2; 1.
Transcription regulation; Zinc-finger.
DOMAIN 415 484 SER-RICH.
ZN_FING 546 RING-TYPE.
DOMAIN 447 461 POLY-SER.
SEQUENCE 600 AA; 66470 MW; E682995:
                                                                                                                                                                                                                                       Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                        between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FUNCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Opposing developmental functions of positive and coregulators of LIM homeodomain factors."; Submitted (JUN-1998) to the EMBL/GenBank/DDBJ data
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RN12_MOUSE
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                                                                                                                                                                                                                                                                                                                                                                                                                                               DEACETYLASE COREPRESSOR COMPLEX.
SUBUNIT: ASSOCIATES WITH LIM/HOMEOBOX
SIMILARITY: CONTAINS 1 RING-TYPE ZINC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: ACTS AS A NEGATIVE CC
TRANSCRIPTION FACTORS. VIA THE
                 LSLCTICSDFFDHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCR 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CPTCRQKV 137
LKTCSVCITEYTEGDKLRKLPCSHEFHVHCIDRWL--SENSTCPICR
                                                                        Similarity
                                                          Conservative
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29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Created)
                                                                       . 88;
                                                         10;
                                                       Score 89; DB
Pred. No. 0.00
LO; Mismatches
                                                                                                                                POLY-SER.
E68299530126E41D
                                                                                                                                                               RING-TYPE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CO-REGULATOR FOR LIM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     600
                                                                        DB 1;
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                                                                                   Length 600
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                                                         Indels
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Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         finger protein)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HOMEODOMAIN
SIN3A/HISTONE
                                                       2
                                                       Gaps
                                                         1;
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RESULT

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Qy

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Conservative

9;

Indels

4;

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LCTICSDFFDHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV

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CELC_HUMAN
IID CELC_HUMAN
AC QOULVB
AC QOULVB
AC QOULVB
DT 16-OCT
RN NCDLIN
RA KIM M.
[1]
RN [1]
RN [2]
RN ESDUEN
RX MEDLLIN
RA Keane
Pennin
RA COOSe
RT ONCOSe
CC -!- FU
CC -!- FU
CC -!- FU
CC -!- FU
CC OT SEE
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CC OT SEE
CC OT 
  Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                   Pfam; PF02262; Cbl_N; 1.
Pfam; PF02761; Cbl_N2; 1.
Pfam; PF02762; Cbl_N3; 1.
Pfam; PF00099; zf-C3HC4; 1.
SMART; SM00184; RING; 1.
SMART; SM00252; SH2; 1.
PR0SITE; PS00518; ZF_RING_1; 1.
PR0SITE; PS05089; ZF_RING_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Penninger J., Lipkowitz S.;

"cbl-3: a new mammalian cbl family protein.";

Oncogene 18:3365-3375(1999).

-i- FUNCTION: REGULATOR OF EGFR MEDIATED SIGNAL TRANSDUCTION.

-i- SUBUNIT: INTERACTS WITH A RESTRICTED RANGE OF SH3 DOMAIN PROTEI

-i- SUBCELLULAR LOCATION: Nuclear (Potential).

-i- ALTERNATIVE PRODUCTS: 2 ISOFORMS; A LONG FORM (SHOWN HERE) AND SHORT FORM; ARE PRODUCED BY ALTERNATIVE SPLICING.

-i- TISSUE SPECIFICITY: UBIQUITOUS.

-i- TISSUE SPECIFICITY: UBIQUITOUS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OSULV8: 09Y5Z3; 09Y5Z2;
16-0CT-2001 (Rel. 40, Created)
16-0CT-2001 (Rel. 40, Last sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Signal transduction protein CBL-C CBLC OR CBL3.
                                                                                                                          CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Pancreatic adenocarcinoma; MEDLINE=99289203; PubMed=10362357;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A. (LONG FORM). MEDLINE=20035821; PubMed=10571044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).

Mortazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or send
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kim M., Tezuka T., Suzuki Y., Sugano S., Hirai M., "Molecular cloning and characterization of a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the European Bioinformatics Institute. Thuse by non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -!- PTM: PHOSPHORYLATED ON TYROSINES BY EGFR.
-!- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Keane M.M., Ettenberg S.A., Nau M.M., Banerjee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. (LONG AND SHORT FORMS)
                                                                                                                                                                          VARSPLIC
                                                                                                                                                                                                 ZN_FING
                                                                                                                                                                                                                          DOMAIN
                                                                                                                                                                                                                                                 Nuclear
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AF117646;
AF117647;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P22681; 1B47
     18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 s requires a license agreement (See http://www.isb-sib.ch/announce/an email to license@isb-sib.ch).
                                                                                                                                                                                                                     protein; Zinc-finger; Phosphorylation; Alternative 89 92 NUCLEAR LOCALIZATION SIGNAL (F
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351
261
234
474
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390
306
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52468
                         30.0%;
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                                                                                                                            MW.
Score 88; DB
Pred. No. 0.00
9; Mismatches
                                                                                                                          RING-TYPE.
MISSING (IN SHORT ISOFORM).
T -> N (IN REF. 2).
; 91013DDF12828242 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Craniata; Vertebrata; Euteleostomi;
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(SH3-binding
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                                                  Length, 474;
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(POTENTIAL)
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RESULT

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MEDLINE-99438124; PubMed=10508479;
Scanlan M.J., Gordan J.D., Williamson
Jongeneal V., Gure A.O., Jager D., Jagold L.J.;
                          Pfam;
                                                                                                                                                                                                                       EMBL;
                                                                                                                                                                                                                                                                use by non-profit institutions as long modified and this statement is not removed. entitles requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                                                               EMBL;
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Ostendorff H.P., Bossenz I
Jenkins N.A., Lichter P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RN12_HUMAN STANDARD; PRT; 624 AA.

QNVW2; QSY598;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
RING finger protein 12 (LIM domain interacting RING finger protein)
(RING finger LIM domain-binding protein) (R-LIM) (NY-REN-43 antigen).
                                                 Pfam;
                                                                       Pfam;
                                                                                                                                                                                                                                                                                                                                                       the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell
                                                                                                                                  InterPro; IPR003837; Glu-tRNAGln.
InterPro; IPR002190; MAGE.
                                                                                                                                                                                                                                                                                                                                                                              between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jenkins N.A., Lichter P., Bach I.; "Functional characterization of the gene encoding RLIM, corepressor of LIM homeodomain transcription factors."; Genomics 69:120-130(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                InterPro; IPR001965;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNF12 OR RLIM.
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                                                                                                                                                                                                                                                                                                                          s SWISS-PROT entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EM European Bioinformatics Institute. There are no restriby non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                            POSITION 134 AND 142.
                                                                                                                                                                        AJ271670; CAC14228.1; -- AF155109; AAD42875.1; ALT_FRAME AK001334; BAA91632.1; --
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            carcinoma
                                                                 PF02686; Glu-tRNAGln;
                                                                                         IPR001841;
                                            MAGE;
                             PHD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      83:456-464(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PubMed=11013082;
                                                                   ; Znf_ring.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody
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                                                                                                                                                                                                                                                                             (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Α.
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E., Knuth
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Hypothetical 47.6 k
C16C10.5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (NOV-1994) to the EMBL/GenBank/DDBJ databases.
-!- SUBCELLULAR LOCATION: Integral membrane protein (Potential).
-!- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
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InterPro; IPR001841; Znf_ring.
Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                       the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Caenorhabditis elegans.
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea.
Rhabditidae; Peloderinae; Caenorhabditis.
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DOMAIN
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8; ZF_RING_1;
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35, Last sequence update)
40, Last annotation update)
kDa protein C16Cl0.5 in chromosome
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6AB33FC9D026C461 CRC64;
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16-OCT-2001
43 kDa recep
                                                                                                         use
                                                                                                                                                          This
                                                                                                                                                                                                                                                                                                                                                                                                  Musil L.S., Carr C., Cohen J.B., Me "Acetylcholine receptor-associated bound myristate.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Frail D.E., mdd J., Shah V., Carr C., Cohen J.B., Mer "cDNAs for the postsynaptic 43-Kba protein of Torpedo encode two proteins with different carboxyl termini."; Proc. Natl. Acad. Sci. U.S.A. 84:6302-6306(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Torpedo californica (Pacific electric ray).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Cl
Elasmobranchii; Squalaa; Hypnosqualaa; Pristiorajea; I
Torpediniformes; Torpedinoidei; Torpedinidae; Torpedo.
EMBL; J02952; AAA49282.1;
EMBL; J02953; AAA49283.1;
PIR; A28009; A28009.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=88107644; PubMed=3427060;
Carr C., McCourt D., Cohen J.B.;
"The 43-kilodalton protein of Torpedo nicotinic postsynaptic
membranes: purification and determination of primary structure.
Biochemistry 26:7090-7102(1987).
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RAPS_TORCA
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43 kDa receptor-associated protein of the synapse (Acetylcholine receptor-associated 43 kDa protein)
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01-MAR-1989
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                                                                                                                                                                                                                                                                                                              Cell Biol. 107:1113-1121(1988).

- FUNCTION: THOUGHT TO PLAY SOME ROLE IN ANCHORING OR STABILIZIN THE NICOTINIC ACETYLCHOLINE RECEPTOR AT SYNAPTIC SITES, IT MAY LINK THE RECEPTOR TO THE UNDERLYING POSTSYNAPTIC CYTOSKELETON, POSSIBLY BY DIRECT ASSOCIATION WITH ACTIN OR SPECTRIN.

- SUBCELLULAR LOCATION: CYTOPLASMIC SURFACE OF POSTSYNAPTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ω
                                                           s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - Buropean Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way ified and this statement is not removed. Usage by and for commercial ities requires a license agreement (See http://www.isb-sib.ch/announce/send an email to license@isb-sib.ch).
                                                                                                                                                                                   ALTERNATIVE PRODUCTS: THERE ARE AT LEAST TWO DISTINCT EXPRESSED, WHICH DIFFER IN THEIR C-TERMINUS. DOMAIN: A CYSTEINE-RICH REGION HOMOLOGOUS TO PART OF REGULATORY DOMAIN OF PROTEIN KINASE C MAY BE IMPORTAN INTERACTIONS OF THIS PROTEIN WITH THE LIPID BILAYER. SIMILARITY: BELONGS TO THE RAPSYN FAMILY. SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
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Pfam; PF009515; TPR; S.
Pfam; PF00097; zf*C3HC4; 1.
PRINTS; PR00217; POSTSYNAPTIC.
PRODOM; PD012428; POSTSYNAPTIC:
1 SMART; SM00184; RING; 1.
SMART; SM00028; TPR; 4.
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SEQUENCE
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PROSITE; PS00518; ZF_RING_1; FALSE_NEG
PROSITE; PS50008; ZF_RING_2; 1
Synapse; Postsynaptic membrane; Cytoske
                                                                                                  by provirus tagging.";
Cell 65:737-752(1991).
                                                                                                                                           SEQUENCE FROM N.A.
MEDLINE=91249381; PubMed=1904008;
van Lohuitzen M., Verbbeek S., Scheijen B.,
                                                                                                                                                                                                             Haupt Y., Alexander W.S., Barri G., Klinken S.P., Adams J "Novel zinc finger gene implicated as myc collaborator by retrovirally accelerated lymphomagenesis in E mu-myc tran
                                                                                                                                                                                                                                                                                                                                             01-MAY-1992 (Rel.
01-MAY-1992 (Rel.
16-OCT-2001 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                       MOUSE
                                                                                                                      Gulden H., Berns A.;
"Identification of cooperating
                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=91249382; PubMed=1904009;
                                                                                                                                                                                                                                                                              NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                         Polycomb complex BMI1 OR BMI-1.
                                                                                                                                                                                                                                                                                                                                                                                P25916;
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                                                                                                                                                                                                                                                                                                                                                                                            BMI1_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VARSPLIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR001237;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 362
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR001440;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 CTICSDFF-DHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCR
SUBCÉLLULAR LOCATION: Nuclear.
TISSUE SPECTEICITY: DEFECTED IN MOST ORGANS WITH
LEVELS IN THYMUS, HEART, BERLIN AND TESTIS.
DISEASE: COOPERATES WITH THE MYC ONCOGENE TO PROI
                                                     FUNCTION: INVOLVED IN MAINTAINING T
STATE OF GENES: MODIFIES CHROMATIN,
IN 1TS EXPRESSIBILITY.
SUBUNIT: COMPONENT OF THE CHROMATIN
                                             (PCG)
                                                                                                                                                                                        65:753-761(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CGLCGESIGDQNSQLQALPCSHLFHLKCL----QTNGNRGCPNCK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 410
411
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA;
                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -finger;
                                                                                                                                                                                                                                                                                                                                   protein
                                                                                                                                                                                                                                                                                                                                               22,
40,
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195
404
411
361
393
                                                                                                                                                                                                                                                                                         Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    46321 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29.9%;
                                                                                                                                                                                                                                                                                                                                               Last
Last
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                                                                                                                                                                                                                                                                                                                                                                     Created)
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                                                                                                                                                                                                                                                                                                                                     BMI - 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 87.5; In Pred. No. 0.00 9; Mismatches
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PHOSPHORYLATION (
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Y -> T (IN REF. 2
Y -> T (IN REF. 2
Y -> T (IN REF. 2
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                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; Sciurognathi; Muridae;
                                                       CHROMATIN-ASSOCIATED
                                                                                                                       oncogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4D26262679FC9B4D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytoskeleton;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     87.5; DB 1;
No. 0.00095;
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on update)
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                                                                                      THE
                                                                            RENDERING
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                                                                                      TRANSCRIPTIONALLY REPRESSIVE
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                                                                                                                                            Wientjens
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                                                                                                                       mu-myc transgenic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ISOFORM)
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                                                                            IT HERITABLY CHANGED
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                                                       POLYCOMB
                                                                                                                                                                                                                                                                                         Euteleostomi;
; Murinae; Mus
                       HIGH
                                                                                                                                                                                                                                   J.M.;
                       EXPRESSION
                                                       COMPLEX
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                                                                                                                                                                                                                                                                                         Mus.
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PRODUCE

B LYMPHOMAS

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В
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Best Local Similarity 33.3%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                              BMI1_HUMAN STANDARE
P35226; Q96E37;
01-FEB-1994 (Rel. 28, C
01-FEB-1994 (Rel. 28, I
01-MAR-2002 (Rel. 41, I
Polycomb complex protei
                                                                                 Strausberg R.;
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
-i- FUNCTION: INVOLVED IN MAINTAINING THE TRANSCRIPTIONALLY REPRESSIVE
STATE OF GENES. MODIFIES CHROMATIN, RENDERING IT HERITABLY CHANGED
IN ITS EXPRESSIBILITY.
                                                                                                                                                                                                            TISSUE=Erythrocyte;
MEDLINE=94093545; PubMed=8268912;
Alkema M.J., Wiegand J., Raap A.K., Berns A., v
"Characterization and chromosomal localization
                                                                                                                                                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalla; Eutherla; Primates;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; M64067; AAA37299.1; -.
EMBL; M64068; AAA37301.1; ALT_SEQ.
EMBL; M64279; AAA37300.1; -.
PIR; A39523; A39523.
PIR; A39524; A39524.
MGD; MGI:88174; Bmi1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                   <del>----</del>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR001841; Znf_ring.
Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00184; RING; 1.
                                                                      -
                                                                                                                                                    TISSUE-Muscle;
                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                        oncogene BMI-1.";
                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                               HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            w
               SUBCELLULAR LOCATION: Nuclear.
DISEASE: COOPERATES WITH THE MYC
SIMILARITY: CONTAINS 1 RING-TYPE
                                                                    SUBUNIT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MCVLCGGYFIDATTI - - IECLHSFCKTCIVRYLET - - SKYCPICDVQV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LCTICSDFFDHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PS00518; ZF_RING_1; 1.
PS50089; ZF_RING_2; 1.
regulator; Nuclear protein; Transcription regulation;
                                                                                                                                                                                                                                                                                                                                                                 complex protein
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81
249
324 AA;
                                                                      COMPONENT
                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
                                                                                                                                                                                        2:1597-1603(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  57 R
95 N
324 P
36707 MW;
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                                                                      OF THE
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Pred. No. 0.00
L2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RING-TYPE.
NUCLEAR LOCALIZATION
PRO/SER-RICH.
                                                                   CHROMATIN-ASSOCIATED
                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AD7DECDB6B29DCE5 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                         update)
                ONCOGENE TO
ZINC FINGER
                                                                                                                                                                                                                                                                                                                                                                                                                                 326
                                                                                                                                                                                                                                                                                                                                                                             update)
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0.00088;
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                                                                                                                                                                                                                                                                                                           Hominidae;
                                                                                                                                                                                                                  van
n of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 324
                              PRODUCE
                                                                                                                                                                                                               Lohuizen M.;
the human proto-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                   POLYCOMB
                                                                                                                                                                                                                                                                                                                         Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60
                            В
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                            LYMPHOMAS
                                                                    COMPLEX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4
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RESULT 16
2147_MOUSE
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                              Orimo A., Inoue S., Ikeda K., Noji S., Muramatsu M.;
"Molecular cloning, structure, and expression of mouse
responsive finger protein Efp. Co-localization with est
mRNA in target organs.";
J. Biol. Chem. 270:24406-24413(1995).
-!- FUNCTION: MEDIATES ESTROGEN ACTION IN VARIOUS TARGE
-!- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
-!- SIMILARITY: CONTAINS 1 SPRY DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation update)
Zinc finger protein 147 (Estrogen responsive)
ZNF147 OR ZFP147 OR EFP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DOMAIN
CONFLICT
SEQUENCE
                                                          EMBL;
                                                                                                                      the European Bioinformatics Institute. use by non-profit institutions as lamodified and this statement is not remo
                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBI the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is
                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. TISSUE=Uterus, Ova
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Repressor; Zinc-finger; Proto-oncogene.

N_FING 18 57 RING-TYPE.

NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).

DOMAIN 81 95 NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).

DOMAIN 251 326 PRO/SER-RICH.

CONFLICT 265 265 V -> I (IN REF. 2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROSITE; PS00518; ZF_RING_1; 1.
PROSITE; PS50089; ZF_RING_2; 1.
Chromatin regulator; Nuclear protein; Transcription regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR001841; Znf_ring.
Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00.184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; L13689; AAA19873.1; -.
EMBL; BC011652; AAH11652.1; -.
MIM; 164831; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                       entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2147_MOUSE
Q61510;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                           MGD; MGI:102749;
                                                                                                                                                                                                                                                                                                                                         TISSUE=Uterus, Ovary, and Placent MEDLINE=96025835; PubMed=7592654;
                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus (Mouse)
                           InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17 MCVLCGGYFIDATTI - · IECLHSFCKTCIVRYLET - · SKYCPICDVQV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 LCTICSDFFDHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV
                                                          D63902; BAA09941.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
16; Conserv
           :102749; Zfp147.
; IPR001870; Gamn
; IPR003877; SPRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  251
265
326 /
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Chordata;
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                           Gamma_carbxylse
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                                                                                                                         is not removed.
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Pred. No. 0.00088;
2; Mismatches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                       (See
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                                                                                                     http://www.
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                                                                                                                       Usage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 326
                                                                                                                                                                                                                                                TARGET ORGANS
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                                                                                                                                                                                                                                                                                              estrogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein) (Efp)
                                                                                                                                                    restrictions
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for
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for
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                                                                                                                                                                      collaboration
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                                                                                                                      in no way commercial
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InterPro;

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RESULT IN RESULT
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Best L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MKR2_HUMAN STANDARD Q9H000; Q9HRY1; Q9BUY2; 01-MAR-2002 (Rel. 41, C 01-MAR-2002 (Rel. 41, L 01-MAR-2002 (Rel. 41, L Makorin 2 (HSPC070).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.

MEDLINE=21481893; PubMed=11597136;
Gray T.A., Azama K., Whitmore K., Min A., Abe S.,
"Phylogenetic conservation of the makorin-2 gene,
zinc-finger protein, antisense to the rafi proto-o
genomics 77:119-126(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DOMAIN
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SMART; SM00184; RING; 1.
SMART; SM00449; SPRY; 1.
PROSITE; PS00518; ZF_RING_1;
PROSITE; PS50089; ZF_RING_2;
Zinc-finger; Coiled coil.
                                                                                                                                                               between
the Euro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zhang Q.-H., Ye M., Wu X.-Y., Ren S.-X., Zhao M., Zhao C.-J., Fu G., Shen Y., Fan H.-Y., Lu G., Zhong M., Xu X.-R., Han Z.-G., Zhang J.-W., Tao J., Huang Q.-H., Zhou J., Hu G.-X., G. J., Chen S.-J., Chen Z.; "Cloning and functional analysis of cDNAs with open reading frames for 300 previously undefined genes expressed in CD34+ hematopoietic stem/progenitor cells.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HUMAN
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Mammalia; Eutheria;
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  EMBL; AF302084; EMBL; AF277170;
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                                                                                                                                                                                                                                                                                                                                                                           rissue=Eye;
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                                                                                                                                                          SWISS-PROT entry is copyright. It is produced through a collable the Swiss Institute of Bioinformatics and the EMBL outst European Bioinformatics Institute. There are no restrictions
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sapiens (Human).

chordata;
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PF00097; zf-C3HC4; 1.
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                                                                   s requires a license agreement (See http://www.isb-sib.an email to license@isb-sib.ch).
                                                                                                                 non-profit institutions as long and this statement is not removed.
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Kawai J., Shinagawa A., Fukunishi Y., Konno H., Adachi J., Fukuda S., Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S., Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I., Salto T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Salto R., Kasukawa T., Salto R., Kasukawa T., Salto R., Flelschmann W., Gassterland T., Gissi C., King B., Kochiwa H., Flelschmann W., Gassterland T., Gissi C., King B., Kochiwa H., Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G., Shake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F., Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M., Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.
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SMART; SM00356; ZnF_C3H1; 4.
PROSITE; PS00518; ZF_RING_1;
PROSITE; PS50089; ZF_RING_2;
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Mammalia; Eutheria; Rodentia;
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15-JUL-1998 (Rel. 3
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      This
                                     Oliver K., Harris D., Barrell B.G., Rajandream M.A., Woo Submitted (AUG-1997) to the EMBL/GenBank/DDBJ databases.-!- SIMILARITY: TO YEAST YHLO10C AND C.ELEGANS EEED8.9.-!- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
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Schizosaccharomycetales;
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InterPro; IPR001841; Znf_ring.
Pfam; PF00097; zf-C3HC4; 1.
Pfam; PF00642; zf-CCCH; 4.
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                                                                                                                                                                                   NCBI_TaxID=4896;
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SMART; SM00356; ZnF_C3H1
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PS00518; zF_RING_1;
PS50089; zF_RING_2;
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C3H1-TYPE 2.
C3H1-TYPE 3.
MAKORIN-TYPE CYS-HIS.
RING-TYPE.
C3H1-TYPE 4.
C3H1-TYPE 4.
L -> F (IN REF. 2).
M; 5F268E6B9D9A6C9F CRC64;
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Pred. No. 0.0013;
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Best Local S
Matches 18
                                                                                   "mel-18, a Polycomb group-related mammalian transcriptional negative regulator with tunc EMBO J. 14:5672-5678(1995)
- FUNCTION: TRANSCRIPTIONAL REPRESSOR. BIN DNA SEQUENCE S'GACTMAGACT-3'. HAS A TUMOR MAY PLAY A ROLE IN CONTROL OF CELL PROLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfam; PF00097; zf-C3HC4; 1.
pfam; PF02148; zf-UBP; 1.
SMART; SM00184; RING; 1.
SMART; SM00290; ZnF-UBP; 1.
PROSITE; PS00518; ZF-RING_1; FALSE_NEG.
PROSITE; PS00699; ZnF-RING_2; 1.
                          <del>;</del>
                                                                                                                                                                           CHARACTERIZATION. MEDLLINE=96091139; PubMed=8521824;
                                                                                                                                                                                                                 Submitted (OCT-2001)
                                                                                                                                                                                                                              Strausberg
                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
MEDLINE=91060627; PubMed=2246278;
Tagawa M., Sakamoto T., Shizanamoto
                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
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ZNF144 OR ZFP144 OR ZFP-144 OR
                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1991 (Rel. 20, Created)
01-MAY-1992 (Rel. 22, Last sequence u
01-MAR-2002 (Rel. 41, Last annotation
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01-NOV-1991
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                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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InterPro; IPR001607; zf-UBP.
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                                                                                                                                                                                                                                                                                      [to T., Nakamura I., Okitsu A., Imai K.,
'Expression of novel DNA-binding protein
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                                               SUBCELLULAR LOCATION: Nuclear TISSUE SPECIFICITY: EXPRESSED
           DEVELOPMENTAL STAGE: DEVELOPMENTALLY REGUING DISEASE: PROBABLY RELATED TO TUMORGENESIS
                                    IN NEURAL TISSUES
                                                                           CELL DEVELOPMENT
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Okitsu A., Imai
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Pred. No. 0.0017;
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ME18_HUMAN
P35227;
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Taniguchi M., Kanno M.;
"Cloning and chromosome mapping
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                                                                             Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.
-- FUNCTION: TRANSCRIPTIONAL REPRESSOR. BINDS SPECIFICALLY TO TO DNA SEQUENCE 5'GACTNGACT-3'. HAS A TUMOR SUPPRESSOR ACTIVITY MAY PLAY A ROLE IN CONTROL OF CELL PROLIFERATION AND/OR NEUR-
                                                                                                                                                                                                                                      Gene
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                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DOMAIN
                                                                                                                                                                                                                                                                                                                                      MEDLINE=93314969;
                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606
                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 LCTICSDFFDHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                  CELL DEVELOPMENT.
SUBCELLULAR LOCATION: Nuclear
TISSUE SPECIFICITY: DETECTED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation - Buropean Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way field and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A37991; A37991.
MGI:99161; Zfp144
                                                                                                                                                                                                                                      les a DNA-binding
129:249-255(1993)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MCALCGGYFIDATTI--VECLHSFCKTCIVRYLET--NKYCPMCDVQV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pro; IPR001841; Znf_ring.
PF00097; zf-C3HC4; 1.
; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BC016419;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              D90085; BAA14122.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                    Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Rel. 28,
(Rel. 28,
(Rel. 40,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          342
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protein Mel-18
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81
242
237
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Human)
    FOUND
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57
RING-
95
NUCLI
342
PRO/S
237
PHOSS
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                                                                                                                                                                                                                                                                                                                 PubMed=8325509;
1., Hasegawa M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              334
                                                                                                                                                                                                                                                                                                                                                                                                                   Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          37723
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence up
annotation
Zinc finge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
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PHOSPHORYLATION (BY CAPK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PHOSPHORYLATION
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C23031B8B9E30108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RING-TYPE
                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (POTENTIAL)
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  IN ALL
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finger protein
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    AND
                                                                                                                                                                                                                                                                             human Mel-18
TISSUE EXAMINED ND KIDNEY AND LOW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               144).
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                                                                                                                                                                                                                                                                                                                 Т.,
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    LOW EXPRESSION
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                                                                               NEURAL
                                                                                                                                                                                                                                                                                                                   M.C.,
                                                                                                                      THE
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RESULT 22
CBL_HUMAN
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Best L
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SEQUENCE
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ZN_FING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entities requires a license agreement (See http://www.isb-sib.ch/announce/
or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         between the Swiss Institute of Bioinformat
the European Bioinformatics Institute. The
use by non-profit institutions as long a
modified and this statement is not removed.
  1 1 1
                                                                                                                                                                         v-cbl was generated by a large truncation domain and a leucine zipper-like motif."; oncogene 6:653-657(1991).
                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                 Signal
                                                                                                                                                                                                                                                                                                                                          01-AUG-1991 (Rel. 19, Created)
01-AUG-1991 (Rel. 19, Last sequence up
16-OCT-2001 (Rel. 40, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                P22681;
                                                                                                                                                                                                                                                                                                                                                                                          CBL_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pfam; PF00097; zf-C3HC4; SMART; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PIR; JN0717; JN0717. MIM; 600346; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through
                                                                                  Nature 398:84-90(1999).
                                                                                                                                         X-RAY CRYSTALLOGRAPHY (2.2 ANGSTROMS) MEDLINE=99176421; PubMed=10078535;
                                                                                                                                                                                                                          Blake T.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS00518; ZF_RING_1; 1.
PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                    Meng W., Sawasdikosol S., Burakoff S.J., "Structure of the amino-terminal domain obliding site on ZAP-70 kinase.";
                                                                                                                                                                                                             "The sequences of
                                                                                                                                                                                                                                                                                                           Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transcription regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro;
                                                                                                                                                                                                                                     MEDLINE=91232862;
                                                                                                                                                                                                                                                 SEQUENCE FROM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 LCTICSDFFDHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IN LIVER, PANCREAS AND SKELETAL MUSCLE.
SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
SURFACE. ASSOCIATES WITH NCK VIA SUBCELLULAR LOCATION: Nuclear. PTM: PHOSPHORYLATED ON TYROSINE. SIMILARITY: CONTAINS 1 RING-TYPE
                                                                                                                                                                                                                                                                                                                       OR CBL2.
                                                        FUNCTION: PARTICIPATES IN SIGNAL TRANSDUCTION IN HEMATOPOIETIC CELLS. ADAPTOR PROTEIN THAT FUNCTIONS AS A NEGATIVE REGULATOR MANY SIGNALLING PATHWAYS THAT START FROM RECEPTORS AT THE CELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                     MCALCGGYFIDATTI - - VECLHSFCKTCIVRYLET - - NKYCPMCDVQV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BC004858;
                                                                                                                                                                                                                                                                                                                               transduction protein CBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 l Similarity 29.:
14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IPR001841;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           242
344 AA;
                                                                                                                                                                                                      Shapiro M., Morse H.C. 111, Lunces of the human and mouse c-cbl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18
                                                                                                                                                                                                                                                 N.A.
                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH04858.1;
                                                                                                                                                                                                                          PubMed=2030914;
ro M., Morse H.C.
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95
344
37788
                                                                                                                                                                                                                                                                                   Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ΜW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Repressor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 86;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEAR LOCALIZATION PRO/SER-RICH.
; A910BCD4C0CE9927 CF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RING-TYPE
                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                           PRT;
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                                                                                                                                                                                                                                                                                                                                 (Proto-oncogene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                          III, Langdon W.Y.;
                                                                                                                                                                                                                                                                                                                                                        update)
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  ZINC FINGER
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                                                                                                                   Eck M.J.;
of Cbl complexed
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                                    DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA-binding
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Best Local
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Pfam; PF02761; Cbl_N2; 1.

Pfam; PF02762; Cbl_N3; 1.

Pfam; PF00627; UBA; 1.

Pfam; PF00097; zf-C3HC4; 1.

SMART; SM00184; RING; 1.

SMART; SM00168; ZF_RING_1; 1.

PROSITE; PS00518; ZF_RING_2; 1.

PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                                                                                                                          YM81_YEAST STANDARD; PRT; 1562 AA. 004781; Q04029; O1-NOV-1997 (Rel. 35, Created) O1-NOV-1997 (Rel. 35, Last sequence update) 16-OCT-2001 (Rel. 40, Last annotation update) Hypothetical 180.2 kDa protein in FAA4-HOR7 in YMR247C OR YM9408.09C OR YM9920.01C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DOMAIN
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL;
PIR; /
PDB; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWI
between
                         Gentles S., Bowman S., Barrell B.G., Rajandream M.A., W Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases -!- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                              SEQUENCE OF 950-1562 FROM STRAIN-S288C / AB972;
                                                                                                                       SEQUENCE OF 1-956 FROM N.A. STRAIN-S288C / AB972;
                                                                                                                                                                                   Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MIM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                   Submitted
                                                                                                              Hunt
                                                                                                                                                                          Saccharomycetales; Saccharomycetaceae;
                                                                                                                                                                                                                                                                                               YEAST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the
                                                                                                                                                            NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3D-structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Proto-oncogene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR000449;
InterPro; IPR001841;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR003153;
                                                                                                                                                                                                                                                                                                                                                 380
                                                                                                                                                                                                                                                                                                                                              s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A43817;
1B47; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             165360;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  X57110; CAA40393.1;
A43817; A43817.
                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                Bowman S., Barrell
d (MAR-1995) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-APR-99
                                                                                                                                                                                                                                                                                                                                                                                                                                                           906 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       124
381
357
477
689
856
700
                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nuclear protein; Zinc-finger; Phosphorylation;
               STRONG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   127
420
476
688
834
895
                                                                                                                                                                                                                                                                                                                                                                                                          29.0%;
35.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           99646
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    UBA.
               ō
                                                                         N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           MW;
               HUMAN
                                                                                                B.G., Rajandream I
EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                               8;
                                                                                                                                                                                                                                                                                                                                                                                                          Score 85; DB 1;
Pred. No. 0.0041;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           PHOSPHORYLATION.
7D686B050204AD8F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ASP/GLU-RICH (ACIDIC)
UBA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEAR LOCALIZATION SIGNAL (POTENTIAL). RING-TYPE. ASP/GLU-RICH (ACIDIC). PRO-RICH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PHOSPHORYLATION.
                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
               ZNF294
                                                                                                            Rajandream M.A.,
                                                                                                                                                                          Saccharomyces
                                                                                                                                                                                                                        intergenic
                                                                                                                                                                                                                                                                                                                                                                                               19;
                                                                                                  databases
                                                                                                                                                                                                                                                                                                                                                                                                                     Length 906
                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                             Walsh
                                                  Walsh
                                                                                                                                                                                                                        region
                                                                                                             S.V.;
                                                                                                                                                                                                                                                                                                                                                                                              4.
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YBR2_YEAST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query
Best L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                    PIR; S45920; S45920.
SGD; S0000266; YBR062C.
InterPro; IPR001841; Znf_;
Pfam; PF00097; zf-C3HC4; SMART; SMO0184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      YBR2_YEA
P38239;
                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-OCT-1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Hypothetical 13.2 kDa protein in ORC2-TIP1 intergenic
YBR062C OR YBR0528.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hypothetical protein; Zinc-finger ZN_FING 1508 1555 RING SEQUENCE 1562 AA; 180185 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SGD; S0004861; YMR247C.
Interpro; IPR00184; Znf_ring.
SMART; SM00184; RING; 1.
PROSITE; PS00518; ZF_RING_1; FALSE_NEG
PROSITE; PS0089; ZF_RING_2; 1.
UNDERSTRE; PS0089; ZF_RING_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                        entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                        Aljinovic
                                                                                                                                                                                                                                                                                                                                                                                                                                    Saccharomycetales; NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1508 CAICYSIL-HAVDRKLPSKTCPTCKNKFHGACLYKWFRSSGNNTCPLCRSEI 1558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; Z48639; CAA88574.1; -. EMBL; Z48756; CAA88657.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the European Bioinformatics Institute. The use by non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation
                                                                                                                             EMBL;
                                                                                                                                                                                                                                                                                                                      STRAIN-S288C;
                                                                                                                                                                                                                                                                                                                                                             Submitted (AUG-1994)
                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Fungi; Ascomycota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Saccharomyces cerevisiae (Baker's yeast).
                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 CTICSDFFDHSRD-----VAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV
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                                                                                                                             z35931;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                              G., Pohl F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                             CAA85005.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saccharomycetaceae;
                                                                                   2nf_ring
                                                                                                                                                                                                                                                                                                                                                                              3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29.0%;
32.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         180185 MW; 97AC65E881362305 CRC64;
                                                                                                                                                                                                                                                                                                                                                                the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                            Pohl T.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saccharomycotina; Saccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No. 0.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  . 0069
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21;
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PROSITE; PS00518; ZF_RING_1; PROSITE; PS50089; ZF_RING_2;

Hypothetical ZN_FING

. protein;

Zinc-finger

RING-TYPE

FALSE_NEG

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THE RESULT TO THE PROPERTY OF 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-9/331014; FUDDWEUT-3100327,
BUING A.L., Benson D., Howard M.J., Margiotta J.F.;
BUING A.L., Benson D., Howard M.J., Margiotta J.F.;
"Chick ciliary ganglion neurons contain transcripts coding for acetylcholine receptor-associated protein at synapses (rapsyn).";
J. Neurosci. 17:5016-5036(1997).
-I- FUNCTION: THOUGHT TO PLAY SOME ROLE IN ANCHORING OR STABILIZING THE NICOTINIC ACETYLCHOLINE RECEPTOR TO SYNAPTIC SITES. IT MAY LINK THE RECEPTOR TO THE UNDERLYING POSTSYNAPTIC CYTOSKELETON, POSSIBLY BY DIRECT ASSOCIATION WITH ACTIN OR SPECTRIN.
-I- SUBCELLULAR LOCATION: CYTOPLASMIC SURFACE OF POSTSYNAPTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHICK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWI
                                                                                                                                                                                                                                                                                                                   EMBL; AF000138; AAB63149.1; InterPro; IPR001237; Postsynar InterPro; IPR001440; TPR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JUL-1999 (Rel. 38, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
43 kDa receptor-associated protein of the synapse (RAPSYN)
(Acetylcholine receptor-associated 43 kDa protein) (43 kDa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            042393;
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Synapse; Postsynaptic membrane; Cytoskeleton; Myristate; Zinc-finger. Myristate; Zinc-finge 0 By SIMILARITY. ZN_FING 362 402 RING-TYPE.
                                                                                            PROSITE; PS00405; 43_KD_POSTSYNAPTIC; 1
PROSITE; PS00518; ZF_RING_1; FALSE_NEG.
PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                                                                                                            PRINTS; PR00217; POSTSYNAPTIC. ProDom; PD012428; Postsynaptic; SMART; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                       entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                use
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Archosauria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gallus gallus (Chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            postsynaptic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RAPS_CHICK
                                                                                                                                                                          SMART;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified and this statement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             European Bioinformatics Institute. There are no rest
by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DOMAIN: A CYSTEINE-RICH REGION HOMOLOGOUS TO PART OF THE REGULATORY DOMAIN OF PROTEIN KINASE C MAY BE IMPORTANT IN INTERACTIONS OF THIS PROTEIN WITH THE LIPID BILAYER. SIMILARITY: BELONGS TO THE RAPBYN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEMBRANES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTICSDFFDHSRD-----VAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SWISS-PROT entry is copyright. It is produced through a collaboratic sen the Swiss Institute of Bioinformatics and the EMBL outstation
                                                                                                                                                                                                                                                                       Pro; IPR001841; 2
PF00515; TPR; 6.
                                                                                                                                                                          SM00028; TPR; 5.
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20; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Rel. 38,
(Rel. 38,
(Rel. 40,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
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                                                                                                                                                                                                                                                                                              _Znf_ring
                                                                                                                                                                                                                                                                                                                                               Postsynaptic
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38
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Pred.
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; Galliformes;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vertebrata; Euteleostomi;
mes; Phasianidae; Phasianinae;
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es 17;
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SEQUENCE
                                                                                                                                                                                                                                       modified and this statement is not removed entities requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hadano S., Ishida Y., Ikeda J.E.;
"The primary structure and genomic organization of five novel transcripts located close to the Huntington's disease gene on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Mammalia; Eutheria;
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16-OCT-2001
Transcription ZN_FING 1:
                                                                                                                                                                            EMBL; AB000468; BAA19122.1; -. EMBL; U95140; AAC52022.1; -.
                                                                                                                                                                                                                                                                                                       between the Swiss Institute of Bioinformathe European Bioinformatics Institute. The European Bioinformatics Institutions as long
                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genomics
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"Identification and characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chiariotti L.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromosome 4p16.3.";
DNA Res. 5:177-186(1998).
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                                      PROSITE; PS00518; ZF_RING_1; 1.
PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                          InterPro; IPR001841; Znf. Pfam; PF00097; Zf-C3HC4;
                                                                                                                                                        MIM; 602850;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY).
-!- SUBUNIT: INTERACTS WITH GSCL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=98140125; PubMed=9479498;
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Senomics 47:258-265(1998)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 similarity).
SUBCELLULAR LOCATION: Nuclear and cytoplasmic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FUNCTION: ENHANCES STEROID RECEPTOR-MEDIATED TRANSCRIPTIONAL ACTIVATION AS WELL AS ACTIVATING BASAL TRANSCRIPTION (BY
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                                                                                          SM00184; RING;
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  regulation;
32 177
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9; Mismatches
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                        Zinc-finger;
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16-0CT-2001 (Rel. 40, 0
16-0CT-2001 (Rel. 40, 1
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                                                           (RNF4) mapping at 4p16.3."
Genomics 47:258-265(1998).
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Hayashizaki Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=C57BL/6J; TISSUE=Embryo; MEDLINE=21085660; PubMed=11217851;
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[2]
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STRAIN=CD-1; TISSUE=Embryo;
MEDLINE=20284895; PubMed=10822263;
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                                                                                                              "Identification and characterization
                                                                                                                                                                                                                                                                                                                                                                                                                    Nature 409:685-690(2001).
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     ACTIVATION
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                                                                                                                                          A., Bruni C.B.;
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16-0CT-2001
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                                                                                   -I- FUNCTION: ENHANCES STEROID RECEPTOR-MEDIATED TRANSCRIPTIONAL ACTIVATION AS WELL AS ACTIVATING BASAL TRANSCRIPTION.
-I- SUBUNIT: INTERACTS WITH GSCL, ANDROGEN RECEPTOR AND TBP.
-I- SUBCELLULAR LOCATION: Nuclear and cytoplasmic.
-I- TISSUE SPECIFICITY: WIDELY EXPRESSED WITH HIGHEST LEVELS IN
                                                                                                                                                                                                    Mol.
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PROSITE; PS50089; ZF_RING_2; 1.
Transcription regulation; Activator; Zinc-finger; Nuclear protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the European Bioinformatics Institute. use by non-profit institutions as 1 modified and this statement is not rem
                                                                                                                                                                                                                         steroid
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Mammalia;
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                                           -!- SIMILARITY: CONTAINS 1 RING-TYPE
                                                                                                                                                                                                                                             "Identification of a novel RING finger
                                                                                                                                                                                                                                                                    Palvimo J.J
                                                                                                                                                                                                                                                                                       Moilanen A.-M., Poukka H.,
                                                                                                                                                                                                                                                                                                                MEDLINE=98378525; PubMed=9710597;
                                                                                                                                                                                                                                                                                                                                       TISSUE=Testis
                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rattus norvegicus (Rat)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright.
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TISSUE SPECIFICITY: IN THE EMBRYO, EXPRESSED PRIMARILY IN THI
DEVELOPING NERVOUS SYSTEM WITH STRONG EXPRESSION IN THE DORS;
ROOT GANGLIA AND GONADS. UBIQUITOUSLY EXPRESSED IN THE ADULT
DEVELOPMENTAL STAGE: EXPRESSION IS DETECTED FROM EMBRYONIC DI
AND CONTINUES THROUGHOUT DEVELOPMENT AND INTO ADULTHOOD.
                                                                  TESTIS
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SUBUNIT: INTERACTS WITH GSCL, ANDROGEN RECEPTOR
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AK019171;
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U95141; AAC53539.1; -.
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                                                                                                                                                                                                                                                                                         Karvonen U.,
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7; Mismatches
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RESULT
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Best Local :
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P29128;
01-DEC-1992
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HRSE; PLOYNG, LUCK, PICTURE PICTURE PICTURE PICTURE PICTURE PROCESSITE; PRO0184; RING; 1.

SMART; SM00184; RING; 1.

PROSITE; PS00518; ZF_RING_1; 1.

PROSITE; PS50089; ZF_RING_2; 1.

PROSITE; PS50089; ZF_RING_2; 1.
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_HSVBJ
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PROSITE; PS50089; ZF_RING_2; 1.
Transcription regulation; Activator; Zinc-finger; Nuclear ZN_FING 136 181 RING-TYPE.
SEQUENCE 194 AA; 21896 MW; 40C13970FC11DFF2 CRC64;
                                                                                                                                                                                                           EMBL;
                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
MEDLINE-92219360; PubMed-1313901;
Mirth U.V., Fraefel C., Vogt B., Vlcek C.,
"Immediate-early RNA 2.9 and early RNA 2.6
"Immediate-early and encode a putative zir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bovine herpesvirus type 1 (strain Jura)
Viruses; dsDNA viruses, no RNA stage; H
Alphaherpesvirinae; Varicellovirus.
NCBL_TaxID-31518;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR001841; Znf_ri
Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00184; RING; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein.";
                                                                                                                                                                                     PIR; B38209;
                                                                                                                                                                  HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          136 CPICMDGYSEIVQNGRLIVSTECGHVFCSQCLRDSLKNA--NTCPTCRKKI 184
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                                                                                                                                                                                                             AJ004801; CAA06138.1;
                                                                                                                                                                                                                                 M84465; AAA46062.1;
                                                                                                                                                                P28990;
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(Rel. 24, Last sequence update)
(Rel. 40, Last annotation updat
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Pred. No. 0.0013;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C:, Paces V., Schwyzer M.;
2.6 of bovine herpesvirus 1
zinc finger transactivator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     THROUGH PHOSPHORYLATION
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RESULT 30 ICPO_HSVBK
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01-APR-1993
16-OCT-2001
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein.";
J. virol. 66:2763-2772(1992).
-i- PTM: THE STRONGLY ACIDIC REGION
ACTIVATION DOMAIN, POSSIBLY REGU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=92219360; PubMed=1313901; Wirth U.V., Fraefel C., Vogt B., Vlcek C., Paces V., Schwyzer M.; "Immediate-early RNA 2.9 and early RNA 2.6 of bovine herpesvirus
                                                                                                                                                                                                                                     PROSITE; PS00518; ZF_RING_1; 1.
PROSITE; PS50089; ZF_RING_2; 1.
Transcription regulation; Trans-acting DNA-binding; Early protein; Repressor; ZN_FING 13 52 RING-TYPE.
                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; M84464; AAA46061.1; -. PIR; A38209; EDBE22. HSSP; P28990; 1CHC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bovine herpesvirus type 1 (strain K22) Viruses; dsDNA viruses, no RNA stage;
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                                                                                                                                                                                                                                                                                                                                                    Pfam; PF00097; zf-C3HC4; SMART; SM00184; RING; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                between the Swiss Institute of Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -!- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
-!- SIMILARITY: TO OTHER HERPESVIRUSES ICPO PROTEIN.
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                                       1 LSLCTICSDFFDHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV
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LGSCCICLDAITGA - - ARALPCLHAFCLACIRRWLEGRP - - TCPLCKAPV
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19; Conser
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676 1
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(Rel. 25, Last sequence up
(Rel. 40, Last annotation
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67701
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38.0%;
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Pred.
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                                                                                                                                                                                              ASP/GLU-RICH (ACIDIC).
; 9BB0683C9BFCA65D CRC64;
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0.0042;
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                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ņ
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Nuclear protein; z
DOMAIN 116
ZN_FING 373
DOMAIN 477
DOMAIN 931
VARSPLIC 812
VARSPLIC 767
VARSPLIC 771
                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CBLB_HUMAN STANDARD; PRT; 982 AA. 013191; 013192; 013193; 16-OCT-2001 (Rel. 40, Created) 16-OCT-2001 (Rel. 40, Last sequence update) 16-OCT-2001 (Rel. 40, Last annotation update) Signal transduction protein CBL-B (SH3-binding
                                                                                                                          PROSITE; PS00518; ZF_RING_1; 1. PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-95303504; PubMed-7784085; MEDLINE-95303504; PubMed-7784085;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                              MIM; 604491;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).

Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                InterPro; IPR003153;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lipkowitz S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9606;
                                                                                                                                                     erPro; IPR000980; SH2.
erPro; IPR000449; UBA.
erPro; IPR001841; Znf_ring.
m; PF02262; CbL_N; 1.
m; PF02761; CbL_N2; 1.
m; PF02762; CbL_N3; 1.
m; PF02762; CbL_N3; 1.
m; PF00697; UBA; 1.
RT; SM00184; RING; 1.
RT; SM00184; RING; 1.
RT; SM00185; UBA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PTM: PHOSPHORYLATED ON TYROSINES. SIMILARITY: CONTAINS 1 RING-TYPE Z SIMILARITY: CONTAINS 1 UBA DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBCELLULAR LOCATION: Nuclear (Potential).
ALTERNATIVE PRODUCTS: 3 ISOFORMS; A LONG R
TRUNCATED ISOFORM 1 AND TRUNCATED ISOFORM
ALTERNATIVE SPLICING.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE SPECIFICITY: EXPRESSED IN LUNG, AS WELL AS FETAL BRAIN AND LIVER AND H. NOT IN ADULT BRAIN, LIVER, PANCREAS, S.
                                                                                                                                                                                                                                                                                                                                        U26710; AAB09291.1; -. U26711; AAB09292.1; -. U26712; AAB09293.1; -. P22681; 1B47.
                                                                                                              Zinc-finger;
                               119
412
701
970
982
770
       109479
er; Phosphorylation; Alternative splicing.
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
RING-TYPE.
PRO-RICH.
UBA.
MISSING (IN TRUNCATED ISOFORM 1).
DVFD -> TYRI (IN TRUNCATED ISOFORM 2).
MISSING (IN TRUNCATED ISOFORM 2).
MISSING (IN TRUNCATED ISOFORM 2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (Potential).

(Potential).

RMS; A LONG FORM (SHOWN HERE)

RMS; A LONG FORM 2; ARE PRODUCED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mitchell J.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZINC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G, KIDNEY, SPLEEN, AND TESTIS, HEMATOPOIETIC CELL LINES, BUS SALIVARY GLAND, OR SKELETAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FINGER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein CBL-B)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Robbins K.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Euteleostomi;
; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FGR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein with
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RAPS_HUMAN
ID RAPS_HI
AC 013702
AC 01- NOV
DT 16-OCT
DE 43 kDa
DE 643 kDa
DE 71SSUE
RX MEDLIN
RA MEDL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR001237; F
InterPro; IPR001440; I
InterPro; IPR001841; Z
Pfam; PF00515; TPR; 5.
INIT_MET
ZN_FING
LIPID
MOD_RES
MOD_RES
                                                                                                                                                                                                                                                      PRINTS; PR00217; POSTSYNAPTIC.
ProDom; PD012428; Postsynaptic;
SMART; SM00184; RING; 1.
SMART; SM00028; TPR; 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for commodified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Buckel A., Beeson D., James M., Vincent A.; "Cloning of cDNA encoding human rapsyn and mapping locus to chromosome 11p11.2-p11.1."; Genomics 35:613-616(1996).

-!- FUNCTION: THOUGHT TO PLAY SOME ROLE IN ANCHORING.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RAPS_HUMAN Q13702;
                                                                                                                         Myristate;
                                                                                                                                                    Synapse;
                                                                                                                                                                              PROSITE;
                                                                                                                                                                                                    PROSITE; PS00405; 43_KD_POSTSYNAPTIC; 1
PROSITE; PS00518; ZF_RING_1; FALSE_NEG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MIM; 601592;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; 233905; CAA83954.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or send
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entities requires a license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=97001170; PubMed=8812503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Muscle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01.NOV-1997 (Rel. 35, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
43 kDa receptor-associated protein of the synapse (RAPSYN)
(Acetylcholine receptor-associated 43 kDa protein) (43 kDa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1997
16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -!- SIMILARITY: BELONGS TO THE RAPSYN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     postsynaptic protein).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              372 LCKICA---ENDKDVKIEPCGHLMCTSCLTAWQE-SDGQGCPFCRCEI 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 LCTICSDFFDHSRDVAAIHCGHTFHLQCLIQWFETAPSRTCPQCRIQV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: THOUGHT TO PLAY SOME ROLE IN ANCHORING OR STABILIZING THE NICOTINIC ACCETYLCHOLINE RECEPTOR AT SYMAPTIC SITES. IT MAY LINK THE RECEPTOR TO THE UNDERLYING POSTSYNAPTIC CYTOSKELETON, POSSIBLY BY DIRECT ASSOCIATION WITH ACTIN OR SPECTRIN. SUBCELLULAR LOCATION: CYTOPLASMIC SURFACE OF POSTSYNAPTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DOMAIN: A CYSTEINE-RICH REGION HOMOLOGOUS TO PART OF THE REGULATORY DOMAIN OF PROTEIN KINASE C MAY BE IMPORTANT II INTERACTIONS OF THIS PROTEIN WITH THE LIPID BILAYER. SIMILARITY: BELONGS TO THE RAPSYN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEMBRANES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            an email to license@isb-sib.ch).
                                                                                                                                    PS50089; zF_RING_2; 1.
Postsynaptic membrane; Cytoskeleton;
Zinc-finger.

0 0

362 402

1 1

195 195

404 404
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            35, Created)
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Primates;
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                                                                                                                                                                                                                                                                                                                                                                                     Znt_ring
                                                                                                                                                                                                                                                                                                                                                                                                                                        Postsynaptic
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Pred. No. 0.00
8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
BY SIMILARITY.
RING-TYPE.
MYRISTATE (BY SIMILARITY).
PHOSPHORYLATION (POTENTIAL).
PHOSPHORYLATION (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
0.0059;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  В
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                                                                                                                                                 Phosphorylation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             50
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RESULT 33
ABJ_PLEMA 1D A33_PLEMA 2D 00208
DI A33_PLEMA 2D 00208
DI 01-0C
DI 01-0C
DI 16-0C
DI 16-0C
DI 16-0C
DI 16-0C
DI 2100-
OS Pleur 2D 200-
OS Pleur 2D 2D 200-
OS Pleur 2D 
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Best I
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                    Pfam; PF00622; SPRY; 1.

Pfam; PF00643; zf-B_box; 1.

Pfam; PF00097; zf-C3HC4; 1.

PRINTS; PR01406; BBOXZNEINGER.

SMART; SM00502; BBC; 1.

SMART; SM00336; BBOX; 1.

SMART; SM00346; RING; 1.

SMART; SM00449; SPRY; 1.

PROSITE; PS50119; ZF_RING_1; 1.

PROSITE; PS50119; ZF_RING_2; 1.

PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A33_PLEWA
Q02084;
Q1-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for comentitles requires a licens a licens as long as its content is in the statement is not removed.
                                                                                                                                                                                                                                                                                                                                      InterPro; IPR001870; Gamma_carbxylse.
InterPro; IPR003877; SPRY.
InterPro; IPR003878; SPRY.domain.
InterPro; IPR0030315; Znf_bbox.
InterPro; IPR001841; Znf_ring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zinc-binding protein A33.
Pleurodeles waltlii (Iberian ribbed newt).
Eukaryota, Metazoa, Chordata, Craniata, Ve
Amphibia, Batrachia, Caudata, Salamandroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-93154311; PubMed=7679068; Bellini M., Lacroix J.-C., Gall J.G.; Bellini M., Lacroix J.-C., Gall J.G.; Routative zinc-binding protein on lampbrush chromosome loops."; Paputative zinc-binding protein on lampbrush chromosome loops."; EMBO J. 12:107-114(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-OCT-1994 (Rel. 30, 01-OCT-1994 (Rel. 30, 16-OCT-2001 (Rel. 40,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR003649; Bbox_C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; L04190; AAA49614.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DEVELOPMENTAL STAGE: IT FIRST APPEARS ON THE CHROMOSOME LOOPS AND IN THE NUCLEOPLASM OF THE GENETIAL VESICLE (GV). IT IS TRANSMITTED TO THE EGG AT GV BREAKDOWN AND APPEARS IN EMBRYONI NUCLEI AT THE MID-BLASTULA STAGE AND IS FOUND IN MANY BUT NOT ALL NUCLEI AT STILL LATER STAGES OF EMBRYOGENESIS.

SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.

SIMILARITY: CONTAINS 1 BOX-TYPE ZINC FINGER.

SIMILARITY: CONTAINS 1 SPRY DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUBCELLULAR LOCATION: Nuclear.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION: MAY BE A NUCLEAR REGULATORY PROTEIN THAT IS STORED IN THE GERMINAL VESICLE FOR USE DURING EARLY EMBRYOGENESIS AND MAY PLAY A ROLE IN THE SYNTHESIS OR PROCESSING OF PRE-MRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DURING OOGENESIS.
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protein; Developmental protein; Coiled
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Pred. No. 0.00
10; Mismatches
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Q9Y252; Q9UF41;
16-OCT-2001 (Rel
16-OCT-2001 (Rel
16-OCT-2001 (Rel
                                                                                 SMART; SMART; SMARTE; PROSITE;
                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collable between the Swiss Institute of Bioinformatics and the EMBL outsithe European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for con
                                                                                                                                                                                                                                                                                                                                                                                    Blum H., Bauersachs S., Mewes H.-W., Gassenhuber J., Wiemann S.; Submitted (DEC-1999) to the EMBL/GenBank/DDJ databases.
-:- TISSUE SPECIFICITY: WEAKLY EXPRESSED IN PERIPHERAL BLOOD, SPLEEN, PROSTATE, TESTIS AND OVARY.
-!- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Cloning and characterization of RNF6, a mapping to 13q12."; Genomics 58:94-97(1999).
                                                                                                                                                                                               EMBL; AJ010347; CAB40414.1; -. EMBL; AJ010346; CAB40413.1; -. EMBL; AL133621; CAB63747.1; -.
                                                                                                                                                                                                                                                            entities requires a license agreement (See http://www.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=99265977; PubMed=Macdonald D.H.C., Lahiri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Leukocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                  Zinc-finger.
                                                                                                                                                            MIM; 604242;
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                                                                                                                               InterPro; IPR001841; Znf_
Pfam; PF00097; zf-C3HC4;
                                                                                                                                                                              HSSP; P28990; 1CHC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Testis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE OF 320-685 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cross N.C.P
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                                                                                   ZF_RING_2;
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                                                                                                   ZF_RING_1;
    424
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B BOX-TYPE.
COILED COIL.
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Pred. No. 0.0044;
2; Mismatches 16;
POLY-ASP.
RING-TYPE.
; 344584773F2E5EFC CRC64;
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                                                     ARG-RICH
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16-OCT-2001 (Rel. 40, Las
16-OCT-2001 (Rel. 40, Las
2inc finger protein 294.
ZNF294 OR KIAA0714.
                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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094822;
                                                                                                                                                                                                      EMBL; AL163249; CAB90430.1; -. EMBL; AL163248; CAB90429.3; -. EMBL; AB01857; BAA34434.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            - i -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nature
[2]
                                                                                    SMART; SM00184; RING; 1.
PROSITE; PS00518; ZF_RING_1; FALSE_NEG
PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nagase T., Ishikawa K.-I., Suyama M., Kikuno R., Miyajima N., Tanaka A., Kotani H., Nomura N., Ohara O.; "Prediction of the coding sequences of unidentified human genes. The complete sequences of 100 new cDNA clones from brain which complete sequences in vitro.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HUMAN
                                                        Zinc-finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-99087487; PubMed-9872452;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 707-1818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Res. 5:277-286(1998).
SIMILARITY: CONTAINS 1 RING-TYPE ZINC
SIMILARITY: STRONG, TO YEAST YMR247C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ICSVCISDYVTGNKLRQLPCMHEFHIHCIDRWL--SENCTCPICR 673
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                                                                                                                                                                             IPR001841; Znf_ring.
   1818
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   AA;
                                  1814
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   205732
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Pred. No. 0.
                               RING-TYPE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
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   416E3E7D4DACA705 CRC64;
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0.0056;
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                           Trail D.E., McLaughlin L.L., Mudd J., Merlie J.P.;

Refail D.E., McLaughlin L.L., Mudd J., Merlie J.P.;

Tidentification of the mouse muscle 43,000-dalton acetylcholine

Treceptor-associated protein (RAPsyn) by cDNA cloning.";

J. Biol. Chem. 263:15602-15607(1988).

-i- FUNCTION: THOUGHT TO PLAY SOME ROLE IN ANCHORING OR STABILIZING

THE NICOTINIC ACETYLCHOLINE RECEPTOR AT SYNAPTIC SITES. IT MAY

LINK THE RECEPTOR TO THE UNDERLYING POSTSYNAPTIC CYTOSKELETON,

POSSIBLY BY DIRECT ASSOCIATION WITH ACTIN OR SPECTRIN.

-i- SUBCELLULAR LOCATION: CYTOPLASMIC SURFACE OF POSTSYNAPTIC

MEMBRANES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 18
                                           PIR; $04488; $04488.

PIR; A31995; A31995;

MGD; MGI99422; Rapsn

InterPro; IPR001237; Po

InterPro; IPR001440; TP

InterPro; IPR001841; Zn

Pfam; PF00515; TPR; 6.
                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1767
                                                                                                                                                        EMBL; X15788; CAA33789.1;
EMBL; J03962; AAA40030.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-OCT-1989 (Rel. 12, Created)
01-AUG-1990 (Rel. 15, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
43 kDa receptor-associated protein of the synapse (RAPSYN)
(Acetylcholine receptor-associated 43 kDa protein) (43 kDa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RAPS
                            PRINTS; PR00217; POSTSYNAPTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Expression of RNA transcripts for the postsynaptic 43 innervated and denervated rat skeletal muscle."; FEBS Lett. 249:229-233(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          postsynaptic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Froehner S.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=89289985; PubMed=2737281;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Muscle
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                                                                                                                                                                                                                                                                                                                        DOMAIN: A CYSTEINE-RICH REGION HOMOLOGOUS TO PART OF THE REGULATORY DOMAIN OF PROTEIN KINASE C MAY BE IMPORTANT IN INTERACTIONS OF THIS PROTEIN WITH THE LIPID BILAYER. SIMILARITY: BELONGS TO THE RAPSYN FAMILY. SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOUSE
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18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Metazoa;
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Rodentia;
                                                            Znf_ring
                                                                                           Postsynaptic
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Pred. No. 0.014;
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Best Local S
Matches 15
                     Query Match
Best Local
    Matches
                                                                                                                                                                                                                     InterPro; IPR003892; CUE.
InterPro; IPR001841; Znf_ring.
InterPro; IPR001841; Znf_ring.
Pfdam; PF002845; CUE; 1.
Pfam; PF00097; zf-C3HC4; 1.
SMART; SM00546; CUE; 1.
SMART; SM00184; RING; 1.
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YUSS_CAEEL

P90859; P90852;

01-NOV-1997 (Rel. 35, C
01-NOV-1997 (Rel. 35, L
16-OCT-2001 (Rel. 40, L
Hypothetical 64.7 kDa p
F26E4.11.
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INIT_MET
ZN_FING
                                                                                                            Hypothetical | ZN_FING 3: SEQUENCE 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOD_RES
MOD_RES
CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          between
the Euro
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PROSITE; PS00405; 43_CD_POSTSYNAPTIC; 1.

PROSITE; PS00518; ZE_RING_1; FALSE_NEG.

PROSITE; PS50089; ZF_RING_2; 1.

Synapse; Postsynaptic membrane; Cytoskeleton;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-BRISTOL N2;
Lightning J., Baynes C.;
Submitted (CCT-1996) to the EMBL/GenBank/DDBJ databases
Submitted (CCT-1996) to The EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     [1]
SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Caenorhabditis elegans.
Eukaryota; Metazoa; Nem
Rhabditidae; Peloderina
NCBI_TaxID=6239;
                                                                                                                                                                                                                                                                                                                                                           EMBL; Z81070; CAB03009.1; --
EMBL; Z81075; CAB03009.1; JOINED
EMBL; Z81075; CAB03049.1; --
EMBL; Z81070; CAB03049.1; JOINED
EMBL; Z81070; CAB03049.1; JOINED
WORMPEP; F26E4.11; CE09695.
                                                                                                                                                                             PROSITE; PS00518; ZF_RING_1; FALSE_NEG PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LIPID
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15; Conser
  l Similarity
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362 402

1 1

195 195

404 404

343 344
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                                                                                                                                        protein;
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  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peloderinae;
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A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nematoda; Chromadorea; rinae; Caenorhabditis.
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                                                                                                                64709 MW;
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                     28.2%;
                                                                                                                                                          Zinc-finger
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Pred. No. 0.00
10; Mismatches
  6,
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WYRISTATE.

PHOSPHORYLATION (POTENTIAL).

PHOSPHORYLATION (POTENTIAL).

DV -> EL (IN REF. 2).
Score 82.5; DB 1;
Pred. No. 0.0054;
6; Mismatches 20;
                                                                                                              RING-TYPE.
; 7C790C238207E49B
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                                                                                                              CRC64;
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                                          Length
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RESULT 38
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                                                                  MKR1_MOUSE
                                                                                                                                                       В
                                                                                    RESULT
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR001395; Aldo_ket_red.
InterPro; IPR000571; Zf-CCCH.
InterPro; IPR001841; Znf_ring.
InterPro; IPR001841; Znf_ring.
InterPro; IPR001842; Zf-C3HC4; 1.
InterPro; IPR001842; Zf-CCCH; 4.
INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; INTERPRO; IN
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01-MAR-2002
01-MAR-2002
01-MAR-2002
                                                                                                                                                                                                                                                                                                                                             ZN_FING
ZN_FING
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gray T.A., Hernandez L., Carey A.H., Schaldach M.A., Smithwick N Rus K., Marshall Graves J.A., Stewart C.L., Nicholls R.D.; "The ancient source of a distinct gene family encoding proteins featuring RING and C(3)H zinc-finger motifs with abundant expressin developing brain and nervous system."; Genomics 66:76-86(2000).
MKR1_MOUSE STANDARD;
Q9QXP6;
Q1-MAR-2002 (Rel. 41, Created)
                                                                                                                                                                                                                                                                                                                                                                                                             ZN_FING
ZN_FING
ZN_FING
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restruct use by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entitles requires a license agreement (See http://www.isb-sor.send an email to license@isb-sib.ch).
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Macropus eugenii (Tammar wallaby).

Macropus eugenii (Tammar wallaby).

Throndata; Craniata; Vertebrata; Euteleus.

Throndontia; Macropodidae; Macropus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A. MEDLINE=20304755; PubMed=10843807;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AF192786; AAF17489.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linc-finger;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         335 CVVCWELLGTSR----RLPCSHQFHDWCLMWWL--AQDSSCPTCRCTI
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                                                                                                                                                       VCGICMEVVYEKANPSERRFGILSNCNHTYCLKCIRKWRSAKQFESKIIKSCPECRI
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17; Conser
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57
86 10
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104
228
259
331
386
52905
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29.8%;
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                                                                                                                                                                                                                                                                                                                                                 MW;
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Pred. No. 0.00
l1; Mismatches
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C3H1-TYPE 2.
C3H1-TYPE 3.
C3H1-TYPE 3.
MAKORIN-TYPE.
RING-TYPE.
C3H1-TYPE 4.
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                                                                                                                                                                                                                                                                                                                                               3H1-TYPE 4.
CB2D9B147433853E CRC64;
                                                                                                                                                                                                                                                           81.5;
No. 0.
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                                                                                                                                                                                                                                                                                 Length 478;
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RESULT 40

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AC Q9UHC7; Q9UEZ

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Best Local
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Mammalia; Eutheria; F
NCBI_TaxID=10090;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "The ancient source of a distinct gene family encoding profeaturing RING and C(3)H zinc-finger motifs with abundant in developing brain and nervous system.";

Genomics 66:76-86(2000).
                                                                                       Homo sapiens (Human Eukaryota; Metazoa; Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZN_FING
ZN_FING
SEQUENCE
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ZN_FING
ZN_FING
DOMAIN
            MEDLINE=20304755;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SMART; SM00184; RING; 1.
SMART; SM00356; ZnF_C3H1; 4.
PROSITE; PS00518; ZF_RING_1;
PROSITE; PS50089; ZF_RING_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF00097; zf-C3HC4; 1
Pfam; PF00642; zf-CCCH; 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MKRN1
                                                                        NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AF192785; AAF17488.1; -. MGD; MGI:1859353; Mkrn1.
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C3H1-TYPE 2.
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C3H1-TYPE 3.
MAKORIN-TYPE CYS-HIS.
RING-TYPE.
C3H1-TYPE 4.
C3H1-TYPE 4.
C6H1-TYPE 4.
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EMBL; AF192789; AAF18979.1; JOIN
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EMBL; AF192797; AAF18979.1; JOIN
EMBL; AF192792; AAF18979.1; JOIN
EMBL; AF192792; AAF18979.1; JOIN
EMBL; AL136812; CAB66746.1; ...
InterPro; IPR001841; Znf_ring.
Pfam; PF00097; Zf-C3HC4; 1.
Pfam; PF00097; Zf-C3HC4; 1.
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"Towards a catalog of human genes and proteins: sequencing and analysis of 500 novel complete protein coding human cDNAs.";
Genome Res. 11:422-435(2001).
-i- TISSUE SPECIFICITY: Ubiquitous.
-i- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGERS.
                                                                                                           CONFLICT
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SMART; SM00356; ZnF_CA1; 4.
SROSITE; PS00518; ZF_RING_1; 1.
PROSITE; PS50089; ZF_RING_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                        modified and this statement is not removentities requires a license agreement (Sor send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=21154917; PubMed=11230166; Wiemann S., Weil B., Wellenreuther R., Ansorge W., Boecher M., Bloecker H., B.
                                                                                                                                                                                                                                  Zinc-finger;
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VCGICMEVVYEKANPSERRFGILSNCNHTYCLKCIRKWRSAKQFESKIIKSCPECRI
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C3H1-TYPE 3.
MAKORIN-TYPE CYS-HIS.
RING-TYPE.
C3H1-TYPE 4.
L -> V
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GR -> AG (IN REF. 1; A
                                               Score 81.5; D
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L1; Mismatches
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